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1000  
P-1-1  
**RAW SEQUENCE LISTING  
ERROR REPORT**

BIO TECHNOLOGY  
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The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/689,159  
Source: 1649  
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Revised 01/29/2002



1647

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
TIME: 08:30:24

Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\05312002\I689159.raw

3 <110> APPLICANT: St. George-Hyslop, Peter  
 4 Rommens, Johanna  
 5 Fraser, Paul  
 7 <120> TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TO ALZHEIMER'S  
 DISEASE  
 9 <130> FILE REFERENCE: 1034/1F808US7  
 11 <140> CURRENT APPLICATION NUMBER: US 09/689,159  
 12 <141> CURRENT FILING DATE: 2000-10-12  
 14 <150> PRIOR APPLICATION NUMBER: US 08/509,359  
 15 <151> PRIOR FILING DATE: 1995-07-31  
 17 <160> NUMBER OF SEQ ID NOS: 183  
 19 <170> SOFTWARE: PatentIn version 3.0  
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 22 <211> LENGTH: 2791  
 23 <212> TYPE: DNA  
 24 <213> ORGANISM: Homo sapiens  
 26 <220> FEATURE:  
 27 <221> NAME/KEY: misc\_feature  
 28 <222> LOCATION: (1)..(2791)  
 29 <223> OTHER INFORMATION: where n may be either a or g or c or t/u, unknown or other  
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 37 tctgggagcc tgcaagtgcac aacagccctt gcggtcctta gacagcttgg cctggaggag 180  
 39 aacacatgaa agaaagaacc tcaagaggct ttgtttctg taaaacagta tttctataca 240  
 41 gttgctccaa tgacagagtt acctgcacccg ttgtcctact tccagaatgc acagatgtct 300  
 43 gaggacaacc acctgagcaa tactgtacgt agccagaatg acaatagaga acggcaggag 360  
 45 cacaacgaca gacggaggct tggccacccct gagccattat ctaatggacg accccagggt 420  
 47 aactcccgcc aggtgggtgg acaagatgag gaagaagatg aggagctgac attgaaatat 480  
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 57 tgctataagg tcatccatgc ctgcttattt atatcatctc tattgttgct gttcttttt 780  
 59 tcattcattt acttggggaa agtgtttaaa acctataacg ttgctgtggc ctacattact 840  
 61 gttgcactcc ttagtctggaa tttgggtgtg gtggaaatga tttccattca ctggaaaggt 900  
 63 ccacttcgac tccagcaggc atatctattt atgattatgt ccctcatggc cctgggttt 960  
 65 atcaagtacc tccctgaatg gactgcgtgg ctcatcttgg ctgtgatttc agtataatgtat 1020  
 67 ttagtggctg ttttgtgtcc gaaagggtcca cttcgtatgc tgggtgaaac agtcaggag 1080  
 69 agaaatgaaa cgtttttcc agctcttatt tactcctcaa caatgggttg gttggtaat 1140  
 71 atggcagaag gagaccggc agctcaaagg agatgttcca aaaattccaa gtataatgca 1200  
 73 gaaaggcacag aaaggggatc acaagacact gttgcagaga atgatgtatgg cgggttcagt 1260  
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 77 cgagctgtc tccaggaact ttccagcagt atccctcgctg gtgaagaccc agagggaaagg 1380

PP. 6, 7-10  
 Does Not Comply  
 Corrected Diskette Needed

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
TIME: 08:30:24

Input Set : A:\Seqlist.txt  
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83	tttgccctta	cattattact	ccttgcatt	ttcaagaaag	cattgcacgc	tcttccaatc	1560
85	tccatcacct	ttgggcttgc	tttctacttt	gccacagatt	atcttgcata	gcctttatg	1620
87	gaccaattag	cattccatca	attttatatac	tagcatattt	gcggtagaa	tcccatggat	1680
89	gttcttctt	tgactataac	caaatactggg	gaggacaag	gtgatttcc	tgtgtccaca	1740
91	tctaaacaaag	tcaagattcc	cggtctggact	tttgcagctt	ccttccaatg	cttcctgacc	1800
93	accttgcact	atggacttt	ggaaggaggt	gcctatagaa	aacgattttg	aacataacttc	1860
95	atcgcagtgg	actgtgtcct	cggtgcagaa	actaccagat	ttgagggacg	aggtaagga	1920
97	gatatgatag	gcccggaaagt	tgctgtgccc	catcagcagc	ttgacgcgtg	gtcacaggac	1980
99	gatttcaactg	acactgcgaa	cttcaggac	tacgggttac	caagaggta	ggtgaagtgg	2040
101	tttaaaccacaa	acggaactct	tcatcttaaa	ctacacgtt	aaaatcaacc	caataattct	2100
103	gtattaactg	aattctgaac	tttcaggag	gtactgtgag	gaagagcagg	caccagcagc	2160
105	agaatgggaa	atggagaggt	ggcagggggt	tccagcttcc	ctttgatttt	ttgctgcaga	2220
107	ctcatccctt	ttaaatgaga	ctgttttcc	cctctcttgc	agtcaagtca	aatatgtaga	2280
109	tgcccttggc	aattcttcctt	ctcaagcact	gacactcatt	accgtctgtg	attgccattt	2340
111	cttcccaagg	ccagtctgaa	cctgagggttgc	ctttatccta	aaagttttaa	cctcagggttc	2400
113	caaattcagt	aaattttgga	aacagtacag	ctattctca	tcaattctct	atcatgttga	2460
115	agtcaaattt	ggattttcca	ccaaattctg	aatttgtaga	catacttgc	cgctcacttg	2520
117	ccccagatgc	ctcctctgtc	ctcattcttc	tctcccacac	aagcagtctt	tttctacagc	2580
119	cagtaaggca	gctctgtcgt	ggtagcagat	ggtcccactt	attctaggtt	cttactcttt	2640
121	gtatgatgaa	aagaatgtgt	tatgaatcgg	tgctgtcagc	cctgctgtca	gaccttcttc	2700
123	cacagcaaataat	gagatgtatg	cccaaagcgg	tagaattaaa	gaagagtaaa	atggctgttg	2760
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128 <210> SEQ ID NO: 2

129 <211> LENGTH: 467

130 <212> TYPE: PRT

131 <213> ORGANISM: Homo sapiens

133 <400> SEQUENCE: 2

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139					20				25			30				
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142					35			40			45					
144	Pro	Leu	Ser	Asn	Gly	Arg	Pro	Gln	Gly	Asn	Ser	Arg	Gln	Val	Val	Glu
145					50			55			60					
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148					65			70			75			80		
150	His	Val	Ile	Met	Leu	Phe	Val	Pro	Val	Thr	Leu	Cys	Met	Val	Val	Val
151						85			90			95				
153	Val	Ala	Thr	Ile	Lys	Ser	Val	Ser	Phe	Tyr	Thr	Arg	Lys	Asp	Gly	Gln
154						100			105			110				
156	Leu	Ile	Tyr	Thr	Pro	Phe	Thr	Glu	Asp	Thr	Glu	Thr	Val	Gly	Gln	Arg
157						115			120			125				
159	Ala	Leu	His	Ser	Ile	Leu	Asn	Ala	Ala	Ile	Met	Ile	Ser	Val	Ile	Val
160						130			135			140				
162	Val	Met	Thr	Ile	Leu	Leu	Val	Val	Leu	Tyr	Lys	Tyr	Arg	Cys	Tyr	Lys
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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
TIME: 08:30:24

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Output Set: N:\CRF3\05312002\I689159.raw

165 Val Ile His Ala Trp Leu Ile Ser Ser Leu Leu Leu Leu Phe Phe  
166 165 170 175  
168 Phe Ser Phe Ile Tyr Leu Gly Glu Val Phe Lys Thr Tyr Asn Val Ala  
169 180 185 190  
171 Val Asp Tyr Ile Thr Val Ala Leu Leu Ile Trp Asn Leu Gly Val Val  
172 195 200 205  
174 Gly Met Ile Ser Ile His Trp Lys Gly Pro Leu Arg Leu Gln Gln Ala  
175 210 215 220  
177 Tyr Leu Ile Met Ile Ser Ala Leu Met Ala Leu Val Phe Ile Lys Tyr  
178 225 230 235 240  
180 Leu Pro Glu Trp Thr Ala Trp Leu Ile Leu Ala Val Ile Ser Val Tyr  
181 245 250 255  
183 Asp Leu Val Ala Val Leu Cys Pro Lys Gly Pro Leu Arg Met Leu Val  
184 260 265 270  
186 Glu Thr Ala Gln Glu Arg Asn Glu Thr Leu Phe Pro Ala Leu Ile Tyr  
187 275 280 285  
189 Ser Ser Thr Met Val Trp Leu Val Asn Met Ala Glu Gly Asp Pro Glu  
190 290 295 300  
192 Ala Gln Arg Arg Val Ser Lys Asn Ser Lys Tyr Asn Ala Glu Ser Thr  
193 305 310 315 320  
195 Glu Arg Glu Ser Gln Asp Thr Val Ala Glu Asn Asp Asp Gly Gly Phe  
196 325 330 335  
198 Ser Glu Glu Trp Glu Ala Gln Arg Asp Ser His Leu Gly Pro His Arg  
199 340 345 350  
201 Ser Thr Pro Glu Ser Arg Ala Ala Val Gln Glu Leu Ser Ser Ser Ile  
202 355 360 365  
204 Leu Ala Gly Glu Asp Pro Glu Glu Arg Gly Val Lys Leu Gly Leu Gly  
205 370 375 380  
207 Asp Phe Ile Phe Tyr Ser Val Leu Val Gly Lys Ala Ser Ala Thr Ala  
208 385 390 395 400  
210 Ser Gly Asp Trp Asn Thr Thr Ile Ala Cys Phe Val Ala Ile Leu Ile  
211 405 410 415  
213 Gly Leu Cys Leu Thr Leu Leu Leu Ala Ile Phe Lys Lys Ala Leu  
214 420 425 430  
216 Pro Ala Leu Pro Ile Ser Ile Thr Phe Gly Leu Val Phe Tyr Phe Ala  
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222 Phe Tyr Ile  
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226 <211> LENGTH: 1929  
227 <212> TYPE: DNA  
228 <213> ORGANISM: Mus musculus  
230 <220> FEATURE:  
231 <221> NAME/KEY: misc\_feature  
232 <222> LOCATION: (1)..(1929)  
233 <223> OTHER INFORMATION: where n may be either a or g or c or t/u, unknown or other  
236 <400> SEQUENCE: 3

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
TIME: 08:30:24

Input Set : A:\Seqlist.txt  
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239 agagaaggaa ccaacacaag acagcagccc ttgcaggctct ttaggcagct tggaggagaa 120  
241 cacatgagag aaagaatccc aagagggtttt gttttctttg agaagggtatt tctgtccagc 180  
243 tgctccaatg acagagatac ctgcacccctt gtcctacttc cagaatgccc agatgtctga 240  
245 ggacagccac tccagcagcg coatccggag ccagaatgac agccaagaac ggcagcagca 300  
247 gcatgacagg cagagacttg acaaccctga gccaatatct aatgggcggc cccagagtaa 360  
249 ctcagaacag gtggtggAAC aagatgagga ggaagacgaa gagctgacat taaaatatgg 420  
251 agccaagcat gtcatcatgc tctttgtccc cgtgaccctc tgcattggctcg tcgtcggtgc 480  
253 caccatcaaa tcagtcagct tctatacccg gaaggacggt cagctaactt acacccatt 540  
255 cacagaaagac actgagactg taggccaaag agccctgcac tcgatcctga atgcggccat 600  
257 catgatcagt gtcattgtca ttatgaccat ctcctgggt gtcctgtata aatacaggtg 660  
259 ctacaaggc atccacgcct ggcttattat ttcatctcg ttgttgcgt tcttttttc 720  
261 gttcatttac tttagggaaag tatttaagac ctacaatgtc kccgtggact acgttacagt 780  
263 agcaactccta atctgaaatt ggggtgtggt cgggatgtt gccatccact gaaaaggccc 840  
265 ctttcgactg cagcaggcgt atctcattat gatcagtgc ctcattggccc tggattttat 900  
267 caagtacctc cccgaatgga ccgcattggct catctggct gtgatttcag tatatgattt 960  
269 ggtggctgtt ttatgtccca aaggcccact tcgtatgctg gttgaaacag ctcaggaaag 1020  
271 aaatgagact ctctttccag ctcttatcta ttccctcaaca atgggtgtgt tggtaat 1080  
273 ggctgaagga gacccagaag cccaaaggag ggtaccaag aaccccaagt ataacacaca 1140  
275 aagagcggag agagagacac aggacagtgg ttctggaaac gatgatgggt gttcagtga 1200  
277 ggagtggggag gcccaaagag acagtcaccc ggggcctcat cgctccactc ccgagtcaag 1260  
279 agctgctgtc caggaacttt ctgggagcat tctaacgagt gaagacccgg aggaaagagg 1320  
281 agtaaaaactt ggactggag atttcattttt ctacagtgtt ctgggtggta aggctcagc 1380  
283 aaccgcccagt ggagactgga acacaaccat agcctgcattk gtgcatac tgcattggcct 1440  
285 gtgccttana ttactcctgc tcgcattta caagaaagg tngccagccc ncccatctc 1500  
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291 ttctttgatt atcaaaaaca caaaaacaga gagcaagccc gaggaggaga ctggtgactt 1680  
293 tcctgtgtcc tcagctaaca aaggcaggac tccagctgga cttctgcagc ttccctccga 1740  
295 gtctccctag ccacccgcac tactggactg tggaaaggag cgtctacaga ggaacggttt 1800  
297 ccaacatcca tcgctgcagc agacgggtgtc cctcaatgtc ttgagagaca aggacaagga 1860  
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304 <210> SEQ ID NO: 4

305 <211> LENGTH: 467

306 <212> TYPE: PRT

307 <213> ORGANISM: Mus musculus

309 <220> FEATURE:

310 <221> NAME/KEY: MISC\_FEATURE

311 <222> LOCATION: (1)..(467)

312 <223> OTHER INFORMATION: where X is unknown or other

315 <400> SEQUENCE: 4

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323 Gln Glu Arg Gln Gln His Asp Arg Gln Arg Leu Asp Asn Pro Glu

324 35 40 45

326 Pro Ile Ser Asn Gly Arg Pro Gln Ser Asn Ser Arg Gln Val Val Glu

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
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Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\05312002\I689159.raw

327 50 55 60  
329 Gln Asp Glu Glu Glu Asp Glu Glu Leu Thr Leu Lys Tyr Gly Ala Lys  
330 65 70 75 80  
332 His Val Ile Met Leu Phe Val Pro Val Thr Leu Cys Met Val Val Val  
333 85 90 95  
335 Val Ala Thr Ile Lys Ser Val Ser Phe Tyr Thr Arg Lys Asp Gly Gln  
336 100 105 110  
338 Leu Ile Tyr Thr Pro Phe Thr Glu Asp Thr Glu Thr Val Gly Gln Arg  
339 115 120 125  
341 Ala Leu His Ser Ile Leu Asn Ala Ala Ile Met Ile Ser Val Ile Val  
342 130 135 140  
344 Ile Met Thr Ile Leu Leu Val Val Leu Tyr Lys Tyr Arg Cys Tyr Lys  
345 145 150 155 160  
347 Val Ile His Ala Trp Leu Ile Ile Ser Ser Leu Leu Leu Leu Phe Phe  
348 165 170 175  
350 Phe Ser Phe Ile Tyr Leu Gly Glu Val Phe Lys Thr Tyr Asn Val Xaa  
351 180 185 190  
353 Val Asp Tyr Val Thr Val Ala Leu Leu Ile Trp Asn Trp Gly Val Val  
354 195 200 205  
356 Gly Met Ile Ala Ile His Trp Lys Gly Pro Leu Arg Leu Gln Gln Ala  
357 210 215 220  
359 Tyr Leu Ile Met Ile Ser Ala Leu Met Ala Leu Val Phe Ile Lys Tyr  
360 225 230 235 240  
362 Leu Pro Glu Trp Thr Ala Trp Leu Ile Leu Ala Val Ile Ser Val Tyr  
363 245 250 255  
365 Asp Leu Val Ala Val Leu Cys Pro Lys Gly Pro Leu Arg Met Leu Val  
366 260 265 270  
368 Glu Thr Ala Gln Glu Arg Asn Glu Thr Leu Phe Pro Ala Leu Ile Tyr  
369 275 280 285  
371 Ser Ser Thr Met Val Trp Leu Val Asn Met Ala Glu Gly Asp Pro Glu  
372 290 295 300  
374 Ala Gln Arg Arg Val Pro Lys Asn Pro Lys Tyr Asn Thr Gln Arg Ala  
375 305 310 315 320  
377 Glu Arg Glu Thr Gln Asp Ser Gly Ser Gly Asn Asp Asp Gly Gly Phe  
378 325 330 335  
380 Ser Glu Glu Trp Glu Ala Gln Arg Asp Ser His Leu Gly Pro His Arg  
381 340 345 350  
383 Ser Thr Pro Glu Ser Arg Ala Ala Val Gln Glu Leu Ser Gly Ser Ile  
384 355 360 365  
386 Leu Thr Ser Glu Asp Pro Glu Glu Arg Gly Val Lys Leu Gly Leu Gly  
387 370 375 380  
389 Asp Phe Ile Phe Tyr Ser Val Leu Val Gly Lys Ala Ser Ala Thr Ala  
390 385 390 395 400  
392 Ser Gly Asp Trp Asn Thr Thr Ile Ala Cys Xaa Val Ala Ile Leu Ile  
393 405 410 415  
395 Gly Leu Cys Leu Xaa Leu Leu Leu Ala Ile Tyr Lys Lys Gly Xaa  
396 420 425 430  
398 Pro Ala Xaa Pro Ile Ser Ile Thr Phe Gly Phe Val Phe Xaa Phe Ala  
399 435 440 445

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
TIME: 08:30:25

Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\05312002\I689159.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:3; N Pos. 5,9,1449,1482,1491,1524  
Seq#:4; Xaa Pos. 192,411,421,432,435,446  
Seq#:5; N Pos. 360  
Seq#:6; N Pos. 4,8,23,42,45,60,68,174,182,239,383,488,525,939  
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Seq#:19; N Pos. 204,221,238,241,313,333,335,339,343,364,370,377  
Seq#:20; N Pos. 5,15,26,28,47,70,104,108,122,124,156,182,184,194,203,217  
Seq#:20; N Pos. 225,240,286,315,321,326  
Seq#:21; N Pos. 31,37,41,43,47,71,88,98,101,115,119,128,135,138,140,144,147  
Seq#:21; N Pos. 157,160,163,166,168,169,182,186,205,207,215,224,245,284,312  
Seq#:21; N Pos. 358  
Seq#:22; N Pos. 18,100,134,139,173,237,319,321,351,352,354,359,376,379,384  
Seq#:22; N Pos. 391,401  
Seq#:24; N Pos. 33,60,104,113,141,156,173,182,219,239,285,289,331,337,358  
Seq#:25; N Pos. 105,107,148,211,235,255,264,269,274,281,302,327,336,350,363  
Seq#:25; N Pos. 399,412  
Seq#:26; N Pos. 46,79,110,116,185,191,462,671,699  
Seq#:27; N Pos. 34,197,215,235,273,307,314,333,395,408,412,436,448  
Seq#:29; N Pos. 216,217,228,278,285,295,315  
Seq#:30; N Pos. 292,305,326  
Seq#:31; N Pos. 17,65,81,98,165,172,176,205,223,260,272,296,307,311,316,335  
Seq#:32; N Pos. 210,236,287,339,360,361,362,366,387,389,390,391,394,420,421  
Seq#:32; N Pos. 427,442,460,466

RAW SEQUENCE LISTING ERROR SUMMARY  
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TIME: 08:30:25

Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\05312002\I689159..raw

Seq#:33; N Pos. 57,217,254,304,340  
Seq#:34; N Pos. 60,107,113,117,123,143,149,196,213,217,233,330,344,360,366  
Seq#:34; N Pos. 369,374,382,387,408,410,425  
Seq#:35; N Pos. 22,83,205  
Seq#:36; N Pos. 135,138,145,150,162,180,196,200,243,250,311,319,347,359,366  
Seq#:36; N Pos. 374,393  
Seq#:37; N Pos. 24,52,81,83,105,121,129,132,147,158,183,209,219,226,257,275  
Seq#:37; N Pos. 301,316  
Seq#:38; N Pos. 177,349,374,492,552,558,561,585,643,650,666,675,706,718,732  
Seq#:38; N Pos. 740,746,756  
Seq#:39; N Pos. 19,48,49,52,57,62,77,81,107,111,113,116,123,144,154,175,181  
Seq#:39; N Pos. 182,186,189,190,197,198,209,211,213,216,219,222,234,244,263  
Seq#:39; N Pos. 265,268,274,278,288,294,295,299,303,305,324  
Seq#:40; N Pos. 17,40,65,119,160,177,233,268,297,367,376,381,408  
Seq#:41; N Pos. 34,52,62,71,75,103,110,145,186,215,218,232,243,251  
Seq#:42; N Pos. 20,28,32,61,69,91,95,117,120,131,140,146  
Seq#:43; N Pos. 293,314,322,324,345,352,355,366,369,379,397,466,487,490,498  
Seq#:43; N Pos. 559,570  
Seq#:44; N Pos. 11,32,238,247,296,309,315,329  
Seq#:45; N Pos. 52,133,160,284,371,373,383,389,390,406,419  
Seq#:46; N Pos. 12,88,185,235,269,301,309,315,319,334,364,379,385  
Seq#:47; N Pos. 3,19,22,32,50,91,92,124,166,171,175,192,212,258,278,303,305  
Seq#:47; N Pos. 325,335,354,357,369,375,422,446,471,479,483  
Seq#:48; N Pos. 12,13,70,120,129,166,177,233,234,291,347,348,372,426  
Seq#:49; N Pos. 64,110,117,170,257,306,315,330,336,353,365  
Seq#:50; N Pos. 19,64,163,225,283,301,373,409,440,443,453,471,475,482,484  
Seq#:51; N Pos. 58,59  
Seq#:52; N Pos. 204  
Seq#:53; N Pos. 8,69,125,126,177,265,314,338,361  
Seq#:54; N Pos. 213,225,240,248,260,296,318,330,368,402,494,500,514,518  
Seq#:55; N Pos. 6,16,19,25,28,52,73,109,190,192,206,232,246,248,275,283,292  
Seq#:55; N Pos. 308,338,389,390,391,416,423,502,515  
Seq#:56; N Pos. 40,166,253,377,379,383,402,418,437,451,484  
Seq#:57; N Pos. 7,147,258,272,305,306,358,374,389,402,413,423,424,426,472  
Seq#:58; N Pos. 78,243,289,345,365,380,386,402,408,409,410,411  
Seq#:59; N Pos. 40,253,301,335,361,379,384,397,404,412,418,425,427,436,455  
Seq#:59; N Pos. 461,494,509  
Seq#:60; N Pos. 11,150,241,242,367  
Seq#:61; N Pos. 356,358,363  
Seq#:62; N Pos. 11,116,260,317,325,361,431,439,445,457,476,491,495  
Seq#:63; N Pos. 13,262,319,433,456  
Seq#:64; N Pos. 30,365,368  
Seq#:65; N Pos. 199,223,233,238,256,282,291  
Seq#:66; N Pos. 7,130,178,197,305,332,339,356,380,415,422,433,436,440  
Seq#:67; N Pos. 289,345,356,407,421,428,433,434,441,443,446  
Seq#:68; N Pos. 4,45,57,309,310,311,347,369  
Seq#:69; N Pos. 88,192,253,283,361,366,372,375,386,426,442,478,485  
Seq#:70; N Pos. 390,400  
Seq#:71; N Pos. 45,385,413,416,424

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/689,159

DATE: 05/31/2002  
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Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\05312002\I689159.raw

Seq#:72; N Pos. 60,92,101,138,160,234,255,263,270,284,297,298  
Seq#:73; N Pos. 135,148,157,158,163,169,235,267,277,321,348,354,379,422  
Seq#:74; N Pos. 7,23,36,48,67,445  
Seq#:75; N Pos. 7,389  
Seq#:76; N Pos. 52,94,104,304,325,401,448  
Seq#:77; N Pos. 101,102,132,228,396,419,439,445,467  
Seq#:78; N Pos. 53,54,115  
Seq#:79; N Pos. 49,236,270,434,443,490  
Seq#:80; N Pos. 14,285,407,472  
Seq#:81; N Pos. 5,159,189,190,191,307,319,343,353  
Seq#:82; N Pos. 33,53,72,81,83,87,124,188,191,233,367,401,416,422,424,439  
Seq#:82; N Pos. 460,471,480,492  
Seq#:83; N Pos. 54,75,257,295,333,386,409,411,414,447  
Seq#:84; N Pos. 18,139,157,170,178,230,257,277,363,373,388,437  
Seq#:85; N Pos. 292,312,351  
Seq#:86; N Pos. 199,446,483  
Seq#:87; N Pos. 392,399,439  
Seq#:88; N Pos. 469,487,500  
Seq#:89; N Pos. 49,114,302,319,343,367,470,484  
Seq#:90; N Pos. 12,20,24,124,155,262,293,299,374,392,419,426,431,458,467  
Seq#:90; N Pos. 475,485,486,488,490,491,494  
Seq#:91; N Pos. 56,139,418,438,439,475,477,501  
Seq#:92; N Pos. 109  
Seq#:93; N Pos. 30,479,484  
Seq#:94; N Pos. 53,145,204,260,266,283,305,318,337,339,342,360,364  
Seq#:95; N Pos. 25,26,77,78,142,164,166,179,232,264,277,295,310  
Seq#:96; N Pos. 6,25,33,68,123,131,165,183,201,202,227,267,273,282,304,309  
Seq#:96; N Pos. 310,315,319,326,342,347,362,367,370,371  
Seq#:97; N Pos. 47,149,182,249,250,277,280,340,345  
Seq#:98; N Pos. 25,159,162,165,196  
Seq#:99; N Pos. 282,369,374,421  
Seq#:100; N Pos. 23,39,44,65,68,84,141,142,179,208,246,276,283,308,380,393  
Seq#:100; N Pos. 406,416,424  
Seq#:101; N Pos. 252,409,434  
Seq#:102; N Pos. 84,146,240,335,414  
Seq#:103; N Pos. 144,171,172,217,219,270,315,352,353,358,359,362,369  
Seq#:104; N Pos. 222  
Seq#:105; N Pos. 84,230,269,283,353,356,371,382,390,407,411  
Seq#:106; N Pos. 190,348,421,439  
Seq#:107; N Pos. 145,157,168,180,183,184,189  
Seq#:108; N Pos. 216,419,471  
Seq#:109; N Pos. 94,96,120,122,359,388,432,459  
Seq#:110; N Pos. 2,4,5,6,7,8,9,10,11,12,13,14,15,16,17,22,34,58,67,173  
Seq#:111; N Pos. 3,4,5,6,7,8,9,10,11,16,36,59,71,173,182,190,204  
Seq#:112; N Pos. 1,2,6,8,10,11,12,13,14,15,16,17,20,21,23,43,70,73,98  
Seq#:113; N Pos. 8,11,12,13,14,15,20,21,44  
Seq#:114; N Pos. 7,9,20,28,30,33,37,43,45,46,47,51,53,54,70,96,109,115,116  
Seq#:114; N Pos. 120,127,135,140,153,157,158,168,171,182,183,186,190,193  
Seq#:114; N Pos. 196,198,211,214,217,219,220,221,225,227,230,231,234,239

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RAW SEQUENCE LISTING ERROR SUMMARY  
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Input Set : A:\Seqlist.txt  
Output Set: N:\CRF3\05312002\I689159.raw

Seq#:115; N Pos. 33,39,65  
Seq#:116; N Pos. 2,3,4,5,6,7,8,12,13,15,19  
Seq#:117; N Pos. 1,2,3,4,5,6,7,8,9,12,13,14,15,16,17,22,23,30,34,126  
Seq#:118; N Pos. 2,5,6,7,8,10,11,12,13,14,15,22,27  
Seq#:119; N Pos. 2  
Seq#:120; N Pos. 11,18,55,67,72,132  
Seq#:121; N Pos. 170  
Seq#:122; N Pos. 130,143,152,153,158,164,230,262,272,316,343,349,374,417  
Seq#:123; N Pos. 11,35,155,336,402,435,441,443  
Seq#:124; N Pos. 119,123,164,178,186,212,261,304,318,322,332,348,351,364  
Seq#:125; N Pos. 52,133  
Seq#:135; N Pos. 5,9  
Seq#:136; Xaa Pos. 192,411,421,432,435,446  
Seq#:137; N Pos. 2137,2144,2152,2157,2160,2163,2180,2203,2213,2216,2225  
Seq#:150; N Pos. 12,532,576,592,626,634,643,777,805,808,811,877,909,919,963  
Seq#:150; N Pos. 990,996,1021,1064  
Seq#:151; N Pos. 94,159,172,254,268,276,385,676  
Seq#:152; N Pos. 263,297,302,312,320,346,355,572,843,890,914,931,991,1119  
Seq#:152; N Pos. 1184,1314,1331,1359,1362,1370,1418,1479,1509,1587,1592  
Seq#:152; N Pos. 1598,1601,1612,1652,1668,1704,1711  
Seq#:153; N Pos. 2,26,28,64,65,85,329,605,777,824,833,1031,1104,1151,1218  
Seq#:153; N Pos. 1238,1239,1300,1650,1798,1810,1834,1844  
Seq#:154; N Pos. 121,138,161,165,181,239,295,662,744,747,748,751,898,903  
Seq#:154; N Pos. 1502,1803,1850,1852,1921,1935  
Seq#:156; N Pos. 144,145,157,168,171,172,217,219,270,315,347,356,357,363  
Seq#:156; N Pos. 364,367,372,374,529,536,676,681,755,804,822,916,929,987  
Seq#:156; N Pos. 1004,1014,1067,1094,1116  
Seq#:157; N Pos. 97,122,190,191,192,307,327,448,474,479,495,498,519,522,538  
Seq#:158; N Pos. 8,46,161,170,182,213,234,238,250,291,312,325,337,364,433  
Seq#:158; N Pos. 434,435  
Seq#:159; N Pos. 22,26,44,47,49,68,81,104,107,119,125,131,137,150,184,188  
Seq#:159; N Pos. 237,248,250,252,274,338,339,340,524,525,562,591,629,659  
Seq#:159; N Pos. 666,691,763,776,789,799,807  
Seq#:160; N Pos. 4,8,23,42,45,60,68,174,182,239,383,488,525,939  
Seq#:179; N Pos. 20  
Seq#:181; N Pos. 13,16  
Seq#:182; N Pos. 14

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.  
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"  
or "Unknown". Please explain source of genetic material in <220> to <223>  
section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)  
(Sec.1.823 of new Rules)

Seq# 127, 128, 129, 130, 131, 132, 139, 140, 141, 142, 143, 144, 145, 146, 147, 148, 149, 163  
Seq# 164, 165, 166, 167, 172, 173, 174, 175, 176, 177, 178, 180, 183

(These are errors in the Sequence Listing)

09/29/59

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(sample of global error)

<210> 127  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<400> 127  
tggagactgg aacacaac

see p. 10

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